VL

D I Q M T Q T T S S L S A S L G D R V T gatatccagatgacccagactacatcctcctgtctgcctctctgggagacagagtcacc I S C R A S Q D I S N Y L N W Y Q Q K P attagttgcagggaacattagcaattatttaaactggtatcagcagaaacca D G T V K L L I Y Y T S I L H S G V P S gatggaactgttaaactcctgatctactacacatcaatattacactcaggagtcccatca R F S G S G S G T D Y S L T I S N L E Q aggttcagtggcagtgggtctggaacagattattctctcaccattagcaacctggagcaa E D F A T Y F C Q Q G N T L P W T F G G gaagattttgccactacttacttttgccaacagggtaatacgcttcggtggaCgttcggaacagctggaaatcaaa

VH

FIG. 1

## Nucleotide/residue numbering shown first followed by Kabat Numbering

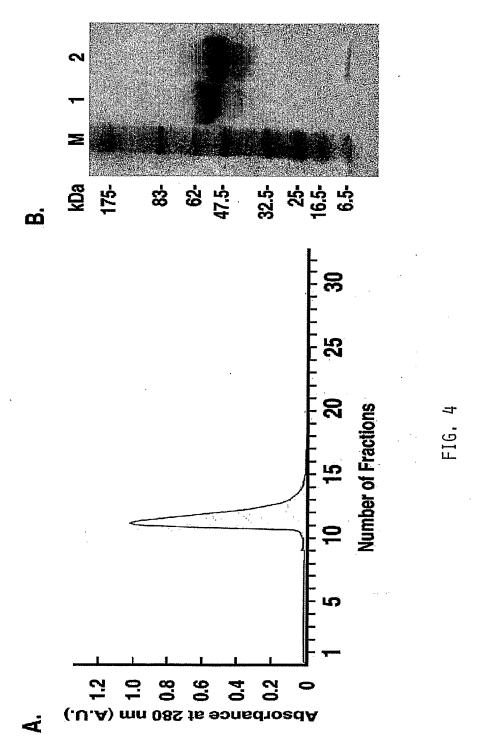
1	0	/ <sup>k</sup>	40 0714 0		
2	1 gat ASP D	49	42 gga GLY G		
3	2 atc ILE I	50	43 act THR T	85	78 ctg LEU L
4	3 cag GLN Q	51	44 gtt VAL V	86	79 gag <b>GLU</b> E
5	4 atg MBT M	52	45 aaa LYS K	87	80 caa <b>GLN</b> Q
6	5 acc THR T	53	46 ctc LEU L	88	81 gaa GLU E
7	6 cag GLN Q	54	47 ctg LEU L	89	82 gat ASP D
		55	48 atc ILE I	90	83 ttt PHE F
8	7 act THR T	56	49 tac TYR Y	91	84 gcc Ala a
9	8 aca THR T	57	50 tac TYR Y	92	85 act THR T
10	9 tcc SER S	<b>5</b> 8	51 aca THR T	93	86 tac TYR Y
11	10 tcc SER S	<b>5</b> 9	52 tca SER S	94	87 ttt PHE F
12	11 ctg LEU L	60	53 ata ILE I	95	88 tgc CYS C
13	12 tct SER S	61	54 tta LEU L	96	89 caa GLN Q
14	13 gcc ALA A	62	55 cac HIS H	97	90 cag GLN Q
15	14 tet ser s	63	56 tca SER S	<b>9</b> 8	91 ggt GLY G
16	15 ctg LEU L	64	57 gga GLY G	99	92 aat ASN N
			58 gtc VAL V		
		65	59 cca PRO P	100	93 acg THR T
17	16 gga GLY G	66		101	94 ctt LEU L
18	17 gac ASP D	67	60 tca SER S	102	95 ccg PRO P
19	18 aga ARG R	68	61 agg ARG R	103	95A
20	19 gtc VAL V	69	62 ttc PHE F	104	95B
21	20 acc THR T	70	63 agt SER S	105	95C
22	21 att ILE I	71	64 ggc GLY G	106 <sup>.</sup>	95D
23	22 agt SER S	72	65 agt SER S	107	95E
24	23 tgc CYS C	73	66 ggg GLY G	108	95F
25	24 agj ARG R	74	67 tct SER S	109	96 tgg TRP W
26	25 gca ALA A	75	68 gga GLY G	110	97 acg THR T
27	26 agt SER S	76	69 aca THR T	111	98 ttc PHE F
28	27 cag GLN Q	<b>7</b> 7	70 gat ASP D	112	99 ggt <b>GLY</b> G
29	27A	<b>7</b> 8	71 tat TYR Y	113	100 gga GLY G
30	27B	<b>7</b> 9	72 tct SER S	114	101 ggc GLY G
31	27C	80	73 ctc LEU L	115	102 acc THR T
32	27D	81	74 acc THR T	116	103 aag LYS K
33	27E	82	75 att ILE I	117	104 ctg LEU L
34	27F	83	76 agc SER S	118	105 gaa GLU E
	28 gac ASP D	84	77 aac ASN N	119	106 atc ILE I
35 36	29 att ILE I	•		120	106A
36	30 age SER S			121	107 aaa LYS K
37	30 age aak a	,		122	108
38	31 aat ASN N			123	109 -
39	32 tat TYR Y				200
40	33 tta LEU L		,		
41	34 aac ASN N				
42	35 tgg TRP W				
43	36 tat TYR Y				
44	37 cag GLN Q				
45	38 cag GLN Q				
46	39 aaa LYS K				
47	40 cca PRO P				
48	41 gat ASP D				

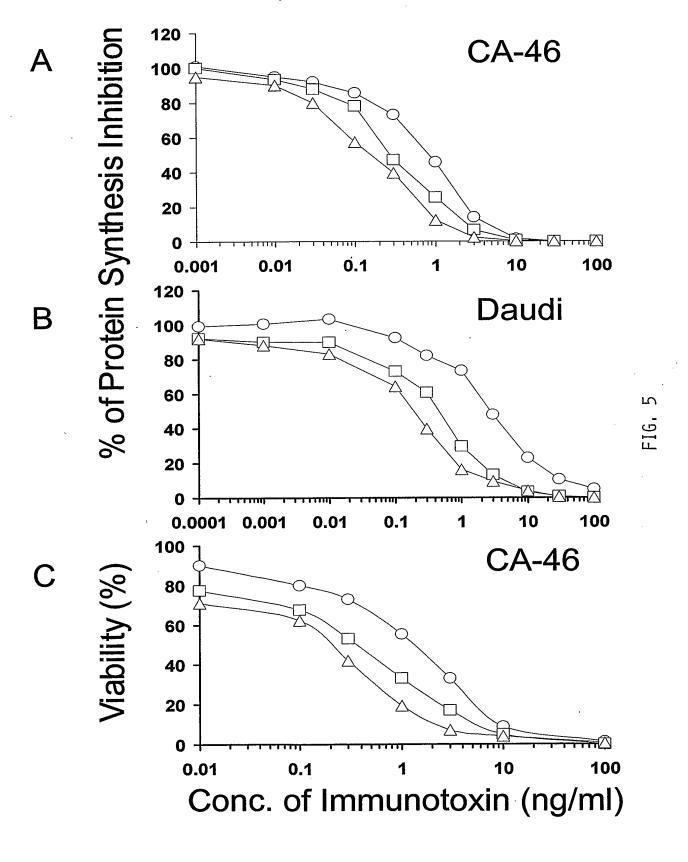
FIG. 2

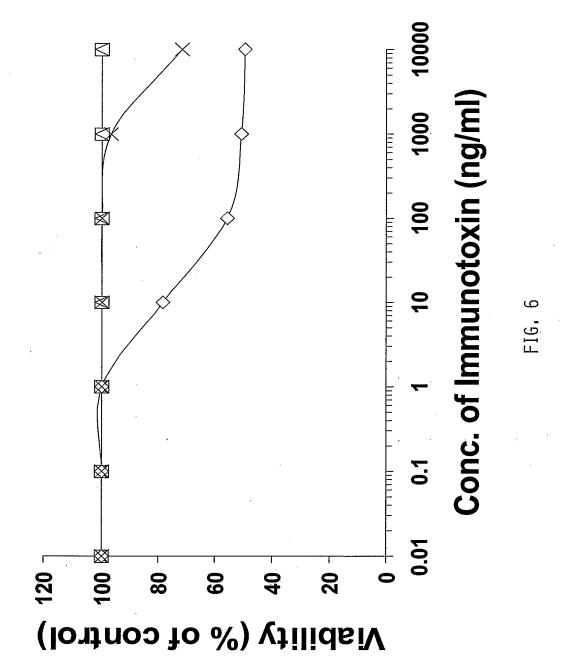
## Nucleotide/residue numbering shown first followed by Kabat Numbering

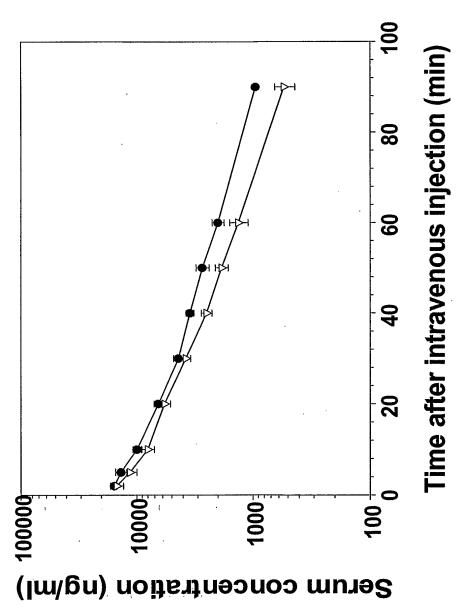
1 0						
1 gaa GLU E 50 47 ggc VAL V 86 80 ctg LEU L 2 gtg VAL V 52 49 gca ALA A 87 81 caa GLN Q 53 50 tac TYR Y 86 82 atg MET M 5 4 ctg LEU L 54 51 att ILE I 89 82A agc SER S 6 5 gtg VAL V 55 52 agt SER S 90 82B agt SER S 7 6 gag GLU G 55 52A agt SER S 90 82B agt SER S 8 8 gg GLY G 55 52A agt SER S 91 82C ctg LEU L 8 8 gg GLY G 55 52B 92 83 aag LYS K 8 8 gg GLY G 55 53 ggt GLY G 94 85 gag GLY G 55 31 gtg GLY G 95 86 gac ASP D 11 10 ggc GLY G 60 54 ggt GLY G 95 86 gac ASP D 12 11 tta LEU L 61 55 ggt GLY G 95 86 gac ASP D 13 12 gtg VAL V 61 55 ggt GLY G 95 86 gac ALA A 13 aag LYS K 62 56 acc THR T 97 88 gcc ALA A 14 13 aag LYS K 63 57 acc THR T 98 89 atg MET M 16 15 gga GLY G 65 55 tat TYR Y 100 91 tac TYR Y 16 15 gga GLY G 65 55 tat TYR Y 100 91 tac TYR Y 16 15 gga GLY G 65 64 58 tac TYR Y 99 90 tat TYR Y 16 15 gga GLY G 65 66 60 cac PRO P 101 92 tgt CYS C 11 tcc SER S 67 61 gac ASP D 102 93 gca ALA A 12 20 tc LEU L 70 64 aag LYS K 105 96 agt SER S 22 21 tcc SER S 71 65 ggc GLY G 106 97 ggc GLY G 12 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 21 tcc SER S 75 69 atc ILE I 110 100A agc SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 100 agt SER S 22 22 tgt CYS C 72 66 cga ARG R 112 100C ggg GLY G 100C agt SER S 111 100B tac TYR Y 100 agt SER S 111 100B tac TYR Y 100 agt SER S 111 100B tac TYR Y 100 agt SER S 111 100B tac TYR Y 100 agt SER S 111 100B tac TYR Y 100 agt SER S 111 100 agt SER S 111 100 agt SER S 112 100 agt SER S 1	1	0			85	70 tag TVD V
3 2 gtg VAL V 52 49 gca ALA A 87 81 caa GIN Q 5 4 ctg LEU L 54 51 att ILE I 89 82 atg MET M 6 5 gtg VAL V 55 52 agt SER S 90 82B agt SER S 8 7 6 gag GLU E 56 52A agt SER S 91 82C ctg LEU L 54 51 att ILE I 89 82A agg SER S 8 8 7 tct SER S 56 52A agt SER S 91 82C ctg LEU L 54 51 att ILE I 89 82A agg SER S 8 8 7 tct SER S 56 52A agt SER S 91 82C ctg LEU L 54 51 9 8 ggg GLY G 58 52C 93 84 tct SER S 10 9 gga GLY G 58 52C 93 84 tct SER S 11 10 ggc GLY G 59 53 ggt GLY G 94 85 gag GLU E 11 tta LEU L 60 54 ggt GLY G 95 86 gac ASP D 12 11 tta LEU L 61 55 ggt GLY G 96 87 aca THR T 13 aag LYS K 62 56 acc THR T 97 88 gcc ALA A 13 aag LYS K 62 56 acc THR T 97 88 gcc ALA A 14 13 aag LYS K 62 56 acc THR T 98 89 atg MET M 15 gga GLY G 65 59 tat TYR Y 100 91 tac TYR Y 16 ggg GLY G 65 59 tat TYR Y 100 91 tac TYR Y 17 16 ggg GLY G 65 59 tat TYR Y 100 91 tac TYR Y 17 100 91 tac TYR Y 17 100 91 tac TYR Y 19 99 90 tat TYR Y 19 100 91 tac TYR Y 19 100 91 t		-		47 Egg TRP W		
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6						
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11 10 ggc GLY G 12 11 tta LEU L 13 12 gtg VAL V 14 13 aag LYS K 15 14 cct PRO P 16 15 gga GLY G 17 16 ggg GLY G 18 17 tcc SER S 17 16 ggg GLY G 18 17 tcc SER S 19 18 ctg LEU L 19 18 ctg LEU L 20 19 aaa LYS K 21 20 tcc LEU L 21 10 tcc SER S 21 10 tcc SER S 22 11 tcc SER S 23 22 tgt CYS C 24 23 gca ALA A 25 24 gcc ALA A 26 acc THR T 27 10 91 tac TYR Y 28 27 tcc PRO P 29 28 gct ALA A 20 29 tcc PRO P 21 20 ctc LEU L 22 21 tcc SER S 25 26 gds GLY G 26 acc THR T 27 10 91 tac TYR Y 28 27 tcc PRO P 29 28 gct ALA A 20 29 tcc PRO P 21 20 ctc LEU L 22 21 tcc SER S 25 26 gds GLY G 26 acc THR T 27 10 92 tcc TYR Y 28 27 tcc PRO P 29 28 gct ALA A 20 29 tcc PRO P 21 20 ctc LEU L 21 20 ctc LEU L 22 21 tcc SER S 23 22 tgt CYS C 24 23 gca ALA A 25 24 gcc ALA A 26 acc THR T 27 10 10 29 3 gca ALA A 27 10 5 ggc GLY G 28 27 tcc PRO P 29 28 gct ALA A 29 28 gct ALA A 29 28 gct ALA A 28 27 6 acc THR T 29 28 gct ALA A 28 27 6 acc THR T 29 28 gct ALA A 29 28 gct ALA A 20 29 tcc PRO P 30 29 tcc PRO P 30 30 dgt SER S 37 35A 39 36 tgg TRP W 40 37 gtt VAL V 41 38 cgc ARG R 42 39 cag GLN Q 43 40 act THR T 44 41 ccg PRO P 45 42 gag GLU E 46 43 aag LYS K 47 44 agg ARG R 48 48 ARG R 49 ARG R 49 ARG R 40 ARG R 40 ARG R 40 ARG R 41 ARG R 41 ARG R 42 ARG R 43 ARG R 44 ARG R 45 CT LEU L 46 ARG R 47 ARG R 48 ARG R 49 ARG R 49 ARG R 40 ARG R 41 ARG R						
12						
13 12 gtg VAL V 62 56 acc THR T 97 88 gcc ALA A 14 13 aag LYS K 62 56 acc THR T 98 89 atg MET M 15 14 cct PRO P 64 58 tac TYR Y 99 90 tat TYR Y 16 15 gga GLY G 65 59 tat TYR Y 100 91 tac TYR Y 17 16 ggg GLY G 66 60 cca PRO P 101 92 tgt CYS C 18 17 tcc SER S 67 61 gac ASP D 102 93 gca ALA A 19 18 ctg LEU L 68 62 act THR T 103 94 aga ARG R 19 18 ctg LEU L 68 62 act THR T 103 94 aga ARG R 20 19 aaa LYS K 69 63 gtg VAL V 104 95 cat HIS H 21 20 ctc LEU L 70 64 aag LYS K 105 96 agt SER S 22 21 tcc SER S 71 65 ggc GLY G 106 97 ggc GLY G 23 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 24 23 gca ALA A 73 67 ttc PHE F 108 99 ggt GLY G 25 24 gcc ALA A 74 68 acc THR T 109 100 agt SER S 26 25 tct SER S 75 69 atc ILE I 110 100A agc SER S 27 26 gga GLY G 76 70 tcc SER S 111 100B tac TYR Y 28 27 ttc PHE F 77 71 aga ARG R 112 100C ggg GLY G 29 28 gct ALA A 78 72 gac ASP D 113 100D gtt VAL V 30 29 ttc PHE F 79 73 aat ASN N 114 100E ttg LEU L 31 30 agt SER S 80 74 gcc ALA A 115 100F 31 30 agt SER S 80 74 gcc ALA A 116 100G 31 30 agt SER S 80 74 gcc ALA A 115 100F 32 31 atc ILE I 81 75 aag LYS K 116 100G 33 32 tat TYR Y 82 76 aac ASN N 117 100H 34 33 gac ASP D 83 77 acc THR T 118 100I 36 35 tct SER S 10 84 78 ctg LEU L 36 35 tct SER S 10 83 77 acc THR T 118 100I 31 30 cgg GLN G 125 105 caa GLN Q 41 30 act THR T 118 100I 32 36 tgg TRP W 124 104 ggc GLY G 125 105 caa GLN Q 41 30 act THR T 126 127 107 act THR T 127 107 act THR T 128 108 ctg LEU L 41 104 ggc GLY G 129 109 gtc VAL V 124 119 109 gtc VAL V 125 119 109 gtc VAL V 126 119 119 119 110 110 act THR T 128 119 119 110 110 act THR T 129 109 gtc VAL V 120 119 119 110 110 act THR T 130 110 act THR T 131 111 tct VAL V 140 140 140 140 140 140 140 140 140 140						
14				55 gge Gui G		
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16		<del>-</del>				
17						
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18		16 ggg GLY G				_
19 19 aaa LYS K 69 63 gtg VAL V 104 95 cat HIS H 21 20 ctc LEU L 70 64 aag LYS K 105 96 agt SER S 22 21 tcc SER S 71 65 ggc GLY G 106 97 ggc GLY G 23 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 24 23 gca ALA A 73 67 ttc PHE F 108 99 ggt GLY G 25 24 gcc ALA A 74 68 acc THR T 109 100 agt SER S 26 25 tct SER S 75 69 atc LLE I 110 100A agc SER S 27 26 gga GLY G 76 70 tcc SER S 111 100B tac TYR Y 28 27 ttc PHE F 77 71 aga ARG R 112 100C ggg GLY G 29 28 gct ALA A 78 72 gac ASP D 113 100D gtt VAL V 30 29 ttc PHE F 79 73 aat ASN N 114 100E ttg LEU L 31 30 agt SER S 80 74 gcc ALA A 115 100F 32 31 atc LLE I 81 75 aag LYS K 116 100G 33 32 tat TYR Y 82 76 aac ASN N 117 100H 34 33 gac ASP D 83 77 acc THR T 118 100I 35 34 atg MET M 84 78 ctg LEU L 119 100J 36 35 tct SER S 37 35A 122 102 tac TYR Y 39 36 tgg TRP W 40 37 gtt VAL V 41 38 cgc ARG R 42 39 cag GLN Q 43 40 act THR T 44 41 ccg PRO P 45 42 gag GLU E 46 43 aag LYS K 47 44 agg ARG R 47 44 agg ARG R 48 46 67 LEU L 48 47 44 agg ARG R				61 gac ASP D		
20 ctc Leu L 70 64 aag Lys K 105 96 agt Ser S 21 20 ctc Leu L 70 64 aag Lys K 106 97 ggc Gly G 22 21 tcc Ser S 71 65 ggc Gly G 106 97 ggc Gly G 23 22 tgt Cys C 72 66 cga Arg R 107 98 tac Tyr Y 24 23 gca Ala A 73 67 ttc Phe F 108 99 ggt Gly G 25 24 gcc Ala A 74 68 acc Thr T 109 100 agt Ser S 26 25 tct Ser S 75 69 atc ILE I 110 100A agc Ser S 27 26 gga Gly G 76 70 tcc Ser S 111 100B tac Tyr Y 28 27 ttc Phe F 77 71 aga Arg R 112 100C ggg Gly G 29 28 gct Ala A 78 72 gac Asp D 113 100D gtt Val V 30 29 ttc Phe F 79 73 aat Asn N 114 100E ttg Leu L 31 30 agt Ser S 80 74 gcc Ala A 115 100F 32 31 atc ILE I 81 75 aag Lys K 116 100G 33 32 tat Tyr Y 82 76 aac Asn N 117 100H 34 33 gac Asp D 83 77 acc Thr T 118 100I 35 34 atg Met M 84 78 ctg Leu L 36 35 tct Ser S 37 35A 38 35B 39 36 tgg Trp W 40 37 gtt Val V 41 38 cgc Arg R 42 39 cag Gln Q 43 40 act Thr T 44 41 ccg Pro P 45 42 gag Glu E 46 43 aag Lys K 47 44 agg Arg R 48 46 67 Leu L						
21 tote SER S 71 65 ggc GLY G 106 97 ggc GLY G 23 22 tgt CYS C 72 66 cga ARG R 107 98 tac TYR Y 24 23 gca ALA A 73 67 ttc PHE F 108 99 ggt GLY G 25 24 gcc ALA A 74 68 acc THR T 109 100 agt SER S 26 25 tct SER S 75 69 atc ILE I 110 100A agc SER S 27 26 gga GLY G 76 70 tcc SER S 111 100B tac TYR Y 28 27 ttc PHE F 77 71 aga ARG R 112 100C ggg GLY G 29 28 gct ALA A 78 72 gac ASP D 113 100D gtt VAL V 30 29 ttc PHE F 79 73 aat ASN N 114 100E ttg LEU L 31 30 agt SER S 80 74 gcc ALA A 115 100F 32 31 atc ILE I 81 75 aag LYS K 116 100G 33 33 gac ASP D 83 77 acc THR T 118 100I 314 33 gac ASP D 83 77 acc THR T 118 100I 135 34 atg MET M 84 78 ctg LEU L 119 100J 136 35 tct SER S 137 35A 122 102 tac TYR Y 123 103 tgg TRP W 124 104 ggc GLY G 126 106 ggg GLY G 127 107 act THR T 128 108 ctg LEU L 140 4 ggc GLY G 127 107 act THR T 144 44 agg ARG R 129 109 gtc VAL V 146 43 aag LYS K 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 147 44 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 131 111 gtc VAL V 148 agg ARG R 148 agg ARG R 131 111 gtc VAL V				63 gtg van v		
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25		22 tgt CYS C		66 CGA ARG R		
26		23 gca ALA A				
26		24 gcc Alia A				
27 26 gga Ghr 6 28 27 ttc PHE F 77 71 aga ARG R 112 100C ggg GLY G 29 28 gct ALA A 78 72 gac ASP D 113 100D gtt VAL V 30 29 ttc PHE F 79 73 aat ASN N 114 100E ttg LEU L 31 30 agt SER S 80 74 gcc ALA A 115 100F 32 31 atc ILE I 81 75 aag LYS K 116 100G 33 32 tat TYR Y 82 76 aac ASN N 117 100H 34 33 gac ASP D 83 77 acc THR T 118 100I 35 34 atg MET M 84 78 ctg LEU L 119 100J 36 35 tct SER S 120 100K ttt PHE F 37 35A 122 102 tac TYR Y 39 36 tgg TRP W 123 103 tgg TRP W 40 37 gtt VAL V 124 104 ggc GLY G 41 38 cgc ARG R 125 105 caa GLN Q 42 39 cag GLN Q 126 106 ggg GLY G 43 40 act THR T 128 108 ctg LEU L 41 ccg PRO P 129 109 gtc VAL V 42 gag GLU E 130 110 act THR T 44 41 ccg PRO P 129 109 gtc VAL V 46 43 aag LYS K 130 110 act THR T 47 44 agg ARG R 131 111 gtc VAL V						
28						
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30		28 gct Ala A				
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35 34 atg MET M 84 78 ctg LEU L 119 100J 120 100K ttt PHE F 121 101 gct ALA A 122 102 tac TYR Y 123 103 tgg TRP W 124 104 ggc GLY G 125 105 caa GLN Q 126 106 ggg GLY G 127 107 act THR T 128 108 ctg LEU L 129 109 gtc VAL V 130 aag LYS K 130 110 act THR T 131 111 gtc VAL V 132 143 44 agg ARG R 131 111 gtc VAL V 132 145 SER S						
35 34 dcg MBY M 36 35 tct SER S 37 35A 121 101 gct ALA A 38 35B 122 102 tac TYR Y 39 36 tgg TRP W 40 37 gtt VAL V 41 38 cgc ARG R 42 39 cag GLN Q 43 40 act THR T 44 41 ccg PRO P 45 42 gag GLU E 46 43 aag LYS K 47 44 agg ARG R 48 120 100K ttt PHE F 121 101 gct ALA A 122 102 tac TYR Y 123 103 tgg TRP W 124 104 ggc GLY G 125 105 caa GLN Q 126 106 ggg GLY G 127 107 act THR T 128 108 ctg LEU L 129 109 gtc VAL V 130 110 act THR T 131 111 gtc VAL V 132 112 tct SER S		33 gac ASP D				
36		of the CED C	04	.0 003		100K ttt PHE F
37 35R 35B 122 102 tac TYR Y 39 36 tgg TRP W 40 37 gtt VAL V 41 38 cgc ARG R 42 39 cag GLN Q 43 40 act THR T 44 41 ccg PRO P 45 42 gag GLU E 46 43 aag LYS K 47 44 agg ARG R 48 45 ctg LEU L 49 ARG R 41 102 tac TYR Y 123 103 tgg TRP W 124 104 ggc GLY G 125 105 caa GLN Q 126 106 ggg GLY G 127 107 act THR T 128 108 ctg LEU L 129 109 gtc VAL V 130 110 act THR T 131 111 gtc VAL V 132 112 tct SER S						101 gct ALA A
38					122	
124 104 ggc GLY G 125 105 caa GLN Q 126 106 ggg GLY G 127 107 act THR T 128 108 ctg LEU L 129 109 gtc VAL V 129 109 gtc VAL V 120 121 122 123 124 125 125 125 125 125 125 125 125 125 125		35B TDD W				103 tgg TRP W
41 38 cgc ARG R 42 39 cag GLN Q 43 40 act THR T 44 41 ccg PRO P 45 42 gag GLU E 46 43 aag LYS K 47 44 agg ARG R 48 125 105 caa GLN Q 126 106 ggg GLY G 127 107 act THR T 128 108 ctg LEU L 129 109 gtc VAL V 130 110 act THR T 131 111 gtc VAL V 132 45 ctg LEU L		30 CGG INF W		•	124	
42 39 cag GLN Q 43 40 act THR T 44 41 ccg PRO P 45 42 gag GLU E 46 43 aag LYS K 47 44 agg ARG R 48 45 ctg LEU L 49 109 gtc VAL V 130 110 act THR T 131 111 gtc VAL V 132 112 tct SER S		37 gcc van v			125	
42 39 cag six v 43 40 act THR T 44 41 ccg PRO P 45 42 gag GLU E 46 43 aag LYS K 47 44 agg ARG R 48 45 ctg LEU L 49 109 gtc VAL V 130 110 act THR T 131 111 gtc VAL V 132 112 tct SER S		38 CGC ARG R				
44 41 ccg PRO P  45 42 gag GLU E  46 43 aag LYS K  47 44 agg ARG R  48 45 ctg LEU L  129 109 gtc VAL V  130 110 act THR T  131 111 gtc VAL V  132 112 tct SER S		40 age mad a			127	107 act THR T
45 42 gag GLU E  46 43 aag LYS K  47 44 agg ARG R  48 45 ctg LEU L  129 109 gtc VAL V  130 110 act THR T  131 111 gtc VAL V  132 112 tct SER S		A1 aca DDO D				
46 43 aag LYS K 47 44 agg ARG R 48 45 ctg LEU L 48 130 110 det IAR I 131 111 gtc VAL V 132 112 tct SER S		42 ded GM E			129	
47 44 agg ARG R 131 111 9tc VAI V		42 gag GEG E				
AS AS CTO LEIL L. 132 112 CCC SER S		44 agg ARG R				
133 113 gca ALA A		45 ctg LEU L				
	***				133	113 gca ALA A

FIG. 3









F16.

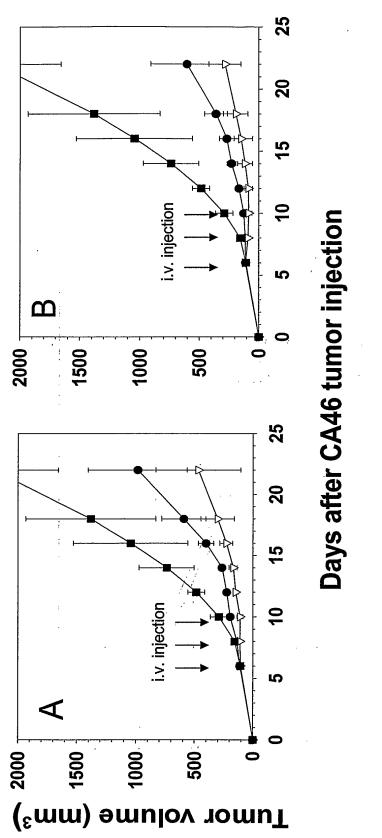


FIG. 8